SEQUENCE LISTING #7

<110> Irwin H. Gelman
Susan G. Jaken

<120> TUMOR SUPPRESSOR GENE

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<212> PRT
<213> Rattus norvegicus

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Glu Pro Gln Glu Val Pro Gln Glu Ala Glu Pro Ala Glu Glu Leu Met
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Lys Ser Arg Glu Met Cys Val Ser Gly Gly Asp His Thr Gln Leu Thr
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Asp Leu Ser Pro Glu Glu Lys Thr Leu Pro Lys His Pro Glu Gly Ile
                85                  90                  95
Val Ser Glu Val Glu Met Leu Ser Ser Gln Glu Arg Ile Lys Val Gln
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Gly Ser Pro Leu Lys Lys Leu Phe Ser Ser Ser Gly Leu Lys Lys Leu
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Ser Gly Lys Lys Gln Lys Gly Lys Arg Gly Gly Gly Gly Asp Glu Glu
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Pro Gly Glu Tyr Gln His Ile His Thr Glu Ser Pro Glu Ser Ala Asp
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Glu Gln Lys Gly Glu Ser Ser Ala Ser Ser Pro Glu Glu Pro Glu Glu
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Thr Thr Cys Leu Glu Lys Gly Pro Leu Glu Ala Pro Arg Met Gly Lys
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Leu Gly Ile Leu Gln Lys Asp Gly Asp Thr Gln Glu Thr Val Arg Arg
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Lys Thr Val Gly Glu Glu Gln Lys Pro Glu Glu Pro Lys Arg Arg Val
                   260                      265                      270
Asp Thr Ser Val Ser Trp Glu Ala Leu Ile Cys Val Gly Ser Ser Lys
                   275                      280                      285
Lys Arg Ala Arg Lys Ala Ser Ser Ser Asp Ile Arg Gly Pro Arg Thr
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Leu Gly Gly Gly Gln Ser Gln Ser Arg Gly Gly Gln Gln Arg Gln Arg
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Ser Arg Thr Asp Ala Val Pro Ala Ser Thr Gln Glu Gln Asp Gln Ala
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Gln Gly Ser Ser Ser Pro Glu Pro Ala Gly Ser Pro Ser Glu Gly Glu
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Gly Val Ser Thr Trp Glu Ser Phe Lys Arg Leu Val Thr Pro Arg Lys
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Lys Ser Lys Ser Lys Leu Glu Glu Lys Glu Ala Gly Arg Thr Leu Val
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Val Gly Ala Gly Cys Pro Leu Arg Ser Asn Arg Val Glu Lys Asn Leu
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Gly Phe Pro Leu Arg Asn Ser Ser Pro Asp Gly Gly Arg Lys Gly Gln
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Met Gly Arg Gln Glu Gln Ala Thr Val Glu Asp Ser Gly Pro Val Glu
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Tyr Asp Ala Val Glu Arg Glu Lys Met Glu Ala Gln Gly Asn Ala Glu
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Leu Pro Ser Cys Trp Gly Cys Val Val Ser Glu Glu Leu Ser Lys Thr
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Val Thr Glu Lys Asp Ile Ile Ala Glu Glu Thr Pro Val Leu Thr Gln
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Thr Leu Pro Glu Gly Lys Asp Ala His Asp Asp Met Val Thr Ser Glu
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Val Asp Phe Thr Ser Glu Ala Val Thr Ala Thr Glu Thr Ser Glu Ala
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Glu Val Leu Ala Ser Glu Lys Glu Lys Asp Val Met Pro Lys Gly Pro
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Gln Ala Val Ala Pro Glu Ser Ser Glu Thr Leu Thr Asp Ser Glu Thr
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Glu Lys Glu Lys Arg Glu Thr Lys Pro Glu Gln Val Ser Glu Glu Gly
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Glu Gln Glu Thr Ala Ala Pro Glu His Glu Arg Asn Tyr Gly Lys Pro
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Pro Leu Gly Pro Glu Ser Gln Ala Glu Ser Ile Pro Ile Ile Val Thr
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Ala Ser Gln Arg Glu Arg Ser Glu Glu Glu Asp Lys Pro Asp Ala Gly
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Pro Asp Ala Asp Gly Lys Glu Ser Thr Ala Ile Asp Lys Val Leu Lys
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catccgaaac cttgacagac agtgagacaa atggaagcac tcccttagca gattcagaca
3480
ctgcagatgg gacacagcaa gatgaaacca ttgacagcca ggacagtaaa gccactgcag
3540
ctgtcaggca gtcacaggtc acagaagaag aggcggctac tgctcagaaa gaggagcctt
3600
cgacactacc taataatgtt ccagcccagg aagaacatgg ggaagaacca ggaagagatg
3660
ttcttgaacc tacacagcaa gagcttactg ctgcagccgt gcccgttctg gcaaagactg
3720
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3780
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3840
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3900
tggaggaggg agagatggaa actgacgttg aaaaggagaa aagggagaca aagccagagc
3960
aagtgagtga agaaggtgag caggaaacag ccgctcctga gcatgaagga acctacggga
4020
agccagtcct gacacttgac atgcccagct cagagagggg gaaggcactg ggaagccttg
4080
gaggaagccc ttctctccca gaccaagaca aagcaggttg catagaggtt caagttcaaa

4140

gcctggacac aacagtcact caaacagcag aagctgtgga aaaggtcata gaaacggttg

4200

tgatttcaga gacaggtgaa agtccagagt gtgtaggtgc acacttatta ccagctgaga

4260

agtcctctgc aacgggtggc cactggactc ttcagcatgc agaggacacg gtacccctgg

4320

ggcctgagtc tcaggcagaa tccatcccaa tcatagtaac tcctgctcct gaaagcaccc

4380

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4440

acaagccaga tgctggtcct gatgctgacg gcaaggagag tacagcaatc gaaaaagtcc

4500

tcaaggctga acctgagatc ctggaacttg agagtaagag caacaagatt gtgctgaacg

4560

tcattcagac agccgttgac cagttcgcac gtacagaaac agcccccgaa actcatgctt

4620

atgattcaca gacccaggtt cctgcatgca ggcttgacag cagggagccc aacagatgct

4680

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4740

aagtcctgac cgttctggag gcatgggctc agcctcggaa atgcttgccg cgcttgcagt

4800

tgaaagcgcc ggtgtcaaag taagcattga gaagctgcct cctcaaccca aagatccaaa

4860

aggagcatgc tgctgatggc cctcagctcc aaagcttagc ccaggcagag gccagtgcct

4920

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4980

atcccccaaa agttgaggtc caggaagaag aaatgtctac caagtcagtc aaagagaaca

5040

aggcccaggc agaagaggac ctgcaggagc caaagggaga cctggcagaa tcctaagatg

5100

ttagttgctc attgtacatc tgtaagacca gaatgtgaaa acaagtcaca gaacaagatg

5160

ctgctgttgg gaccttgaga ccaagatttc agagcccatg agatccagag agcagggccg

5220

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5280

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5340

tatttctaac atttcctcgt ttcaagactg cctttgattt gccccttgat gccgtccgtg

5400

tatttcggat ttaaggtcct gcgttctcaa cctggaacca attctgccat acctagttcc

5460

acttctcaaa ctggagcatc ctcctttatg tatttatatg tatgttttat gtagtcctcc

5520

tcctgtacct attgtatatt tttttctaac gtttaagcac atgctttttg tattatgcaa

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5580
tatataacgg gtgtgcagcc atagcgacgc tttgaaaagc tccaagcctc aactgtaacc
5640
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5700
gcttagatct gtgggcttct agtcctctga aagtggttgt tttcctatgc acagcgagct
5760
cagaaataaa aaccccattt tgaaacatcc aggatgtccc aatattacca tgattttttc
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ccccttttt gctaatccag tccaggttgg aaagaagtct cctctgtgtc agattaagcc
5880
ctgtctctta atgatatgga caaatgagtg tgcctaaggc catgagatgt ttcctaatgc
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agaaggaatc tgttgtacgt tttttgatt gtactcttct atgctggacc gaattcatat
6000
gcagatcgaa gtgagtcctg ttctttacag atggtatttt gatagatact ggagtttgtc
6060
tgtgttatat ctgtgcccct tctttaagaa caatgttgca ttatgttcct ttggataaat
6120
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6160
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<210> 4
<211> 1596
<212> PRT
<213> Rattus norvegicus

<400> 4
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Gly Ser Asp Thr Pro Ser Glu Leu Val Leu Ser Gly His Gly Pro Ala
            20                  25                  30
Ala Glu Ala Ser Gly Ala Ala Gly Asp Pro Ala Asp Ala Asp Pro Ala
            35                  40                  45
Thr Lys Leu Pro Gln Lys Asn Gly Gln Leu Ser Ser Val Asn Gly Val
        50                  55                  60
Ala Glu Gln Gly Asp Val His Val Gln Glu Glu Asn Gln Glu Gly Gln
65                  70                  75                  80
Glu Glu Glu Val Val Asp Glu Asp Val Gly Gln Arg Glu Ser Glu Asp
                85                  90                  95
Val Arg Glu Lys Asp Arg Val Glu Glu Met Ala Ala Asn Ser Thr Ala
            100                 105                 110
Val Glu Asp Ile Thr Lys Asp Gly Gln Glu Glu Thr Ser Glu Ile Ile
        115                 120                 125
Glu Gln Ile Pro Ala Ser Glu Asn Asn Val Glu Glu Met Val Gln Pro
    130                 135                 140
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Ala Glu Ser Gln Ala Asn Asp Val Gly Phe Lys Lys Val Phe Lys Phe
145             150             155             160
Val Gly Phe Lys Phe Thr Val Lys Lys Asp Lys Asn Glu Lys Ser Asp
        165             170             175
Thr Val Gln Leu Leu Thr Val Lys Lys Asp Glu Gly Glu Gly Ala Glu
        180             185             190
Ala Ser Val Gly Ala Gly Asp His Gln Glu Pro Ser Val Glu Thr Ala
        195             200             205
Val Gly Glu Ser Ala Ser Lys Glu Ser Glu Leu Lys Gln Ser Thr Glu
210             215             220
Lys Gln Glu Gly Thr Leu Lys Gln Glu Gln Ser Ser Thr Glu Ile Pro
225             230             235             240
Leu Gln Ala Glu Ser Asp Gln Ala Ala Glu Glu Glu Ala Lys Asp Glu
        245             250             255
Gly Glu Glu Lys Gln Glu Lys Glu Pro Thr Lys Ser Pro Glu Ser Pro
        260             265             270
Ser Ser Pro Val Asn Ser Glu Thr Thr Ser Ser Phe Lys Lys Phe Phe
        275             280             285
Thr His Gly Trp Ala Gly Trp Arg Lys Lys Thr Ser Phe Lys Lys Ser
        290             295             300
Lys Glu Asp Asp Leu Glu Thr Ala Glu Lys Arg Lys Glu Gln Glu Ala
305             310             315             320
Glu Lys Val Asp Glu Glu Glu Lys Glu Lys Thr Glu Pro Ala Ser Glu
        325             330             335
Glu Gln Glu Pro Ala Glu Asp Thr Asp Gln Ala Arg Leu Ser Ala Asp
        340             345             350
Tyr Glu Lys Val Glu Leu Pro Leu Glu Asp Gln Val Gly Asp Leu Glu
        355             360             365
Ala Ser Ser Glu Glu Lys Cys Ala Pro Leu Ala Thr Glu Val Phe Asp
370             375             380
Glu Lys Met Glu Ala His Gln Glu Val Val Ala Glu Val His Val Ser
385             390             395             400
Thr Val Glu Lys Thr Glu Glu Glu Gln Gly Gly Gly Gly Glu Ala Glu
        405             410             415
Gly Gly Val Val Val Glu Gly Thr Gly Glu Ser Leu Pro Pro Glu Lys
        420             425             430
Leu Ala Glu Pro Gln Glu Val Pro Gln Glu Ala Glu Pro Ala Glu Glu
        435             440             445
Leu Met Lys Ser Arg Glu Met Cys Val Ser Gly Gly Asp His Thr Gln
450             455             460
Leu Thr Asp Leu Ser Pro Glu Glu Lys Thr Leu Pro Lys His Pro Glu
465             470             475             480
Gly Ile Val Ser Glu Val Glu Met Leu Ser Ser Gln Glu Arg Ile Lys
        485             490             495
Val Gln Gly Ser Pro Leu Lys Lys Leu Phe Ser Ser Ser Gly Leu Lys
        500             505             510
Lys Leu Ser Gly Lys Lys Gln Lys Gly Lys Arg Gly Gly Gly Gly Asp
        515             520             525
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Glu Glu Pro Gly Glu Tyr Gln His Ile His Thr Glu Ser Pro Glu Ser
530   535   540
Ala Asp Glu Gln Lys Gly Glu Ser Ser Ala Ser Ser Pro Glu Glu Pro
545   550   555   560
Glu Glu Thr Thr Cys Leu Glu Lys Gly Pro Leu Glu Ala Pro Gln Asp
                565   570   575
Gly Glu Ala Glu Glu Gly Thr Thr Ser Asp Gly Glu Lys Lys Arg Glu
            580   585   590
Gly Ile Thr Pro Trp Ala Ser Phe Lys Lys Met Val Thr Pro Lys Lys
            595   600   605
Arg Val Arg Arg Pro Ser Glu Ser Asp Lys Glu Glu Glu Leu Glu Lys
        610   615   620
Val Lys Ser Ala Thr Leu Ser Ser Thr Asp Ser Thr Val Ser Glu Met
625   630   635   640
Gln Asp Glu Val Lys Thr Val Gly Glu Glu Gln Lys Pro Glu Glu Pro
                645   650   655
Lys Arg Arg Val Asp Thr Ser Val Ser Trp Glu Ala Leu Ile Cys Val
                660   665   670
Gly Ser Ser Lys Lys Arg Ala Arg Lys Ala Ser Ser Ser Asp Asp Glu
        675   680   685
Gly Gly Pro Arg Thr Leu Gly Gly Asp Ser His Arg Ala Glu Glu Ala
    690   695   700
Ser Lys Asp Lys Glu Ala Gly Thr Asp Ala Val Pro Ala Ser Thr Gln
705   710   715   720
Glu Gln Asp Gln Ala Gln Gly Ser Ser Ser Pro Glu Pro Ala Gly Ser
            725   730   735
Pro Ser Glu Gly Glu Gly Val Ser Thr Trp Glu Ser Phe Lys Arg Leu
        740   745   750
Val Thr Pro Arg Lys Lys Ser Lys Ser Lys Leu Glu Glu Lys Ala Glu
        755   760   765
Asp Ser Ser Val Glu Gln Leu Ser Thr Glu Ile Glu Pro Ser Arg Glu
770   775   780
Glu Ser Trp Val Ser Ile Lys Lys Phe Ile Pro Gly Arg Arg Lys Lys
785   790   795   800
Arg Ala Asp Gly Lys Gln Glu Gln Ala Thr Val Glu Asp Ser Gly Pro
                805   810   815
Val Glu Ile Asn Glu Asp Asp Pro Asn Val Pro Ala Val Val Pro Leu
            820   825   830
Ser Glu Tyr Asn Ala Val Glu Arg Glu Lys Met Glu Ala Gln Gly Asn
        835   840   845
Thr Glu Leu Pro Gln Leu Leu Gly Ala Val Tyr Val Ser Glu Glu Leu
    850   855   860
Ser Lys Thr Leu Val His Thr Val Ser Val Ala Val Ile Asp Gly Thr
865   870   875   880
Arg Ala Val Thr Ser Val Glu Glu Arg Ser Pro Ser Trp Ile Ser Ala
                885   890   895
Ser Val Thr Glu Pro Leu Glu His Thr Ala Gly Glu Ala Met Pro Pro
            900   905   910

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Val Glu Glu Val Thr Glu Lys Asp Ile Ile Ala Glu Glu Thr Pro Val
        915                 920                 925
Leu Thr Gln Thr Leu Pro Glu Gly Lys Asp Ala His Asp Asp Met Val
        930                 935                 940
Thr Ser Glu Val Asp Phe Thr Ser Glu Ala Val Thr Ala Thr Glu Thr
945                 950                 955                 960
Ser Glu Ala Leu Arg Thr Glu Glu Val Thr Glu Ala Ser Gly Ala Glu
                965                 970                 975
Glu Thr Thr Asp Met Val Ser Ala Val Ser Gln Leu Thr Asp Ser Pro
                980                 985                 990
Asp Thr Thr Glu Glu Ala Thr Pro Val Gln Glu Val Glu Ser Gly Val
        995                 1000                1005
Leu Asp Thr Glu Glu Glu Glu Arg Gln Thr Gln Ala Ile Leu Gln Ala
    1010                1015                 1020
Val Ala Asp Lys Val Lys Glu Glu Ser Gln Val Pro Ala Thr Gln Thr
1025                1030                1035                1040
Val Gln Arg Thr Gly Ser Lys Ala Leu Glu Lys Val Glu Glu Val Glu
                1045                1050                1055
Glu Asp Ser Glu Val Leu Ala Ser Glu Lys Glu Lys Asp Val Met Pro
            1060                1065                1070
Lys Gly Pro Val Gln Glu Ala Gly Ala Glu His Leu Ala Gln Gly Ser
        1075                1080                1085
Glu Thr Gly Gln Ala Thr Pro Glu Ser Leu Glu Val Pro Glu Val Thr
    1090                1095                1100
Ala Asp Val Asp His Val Ala Thr Cys Gln Val Ile Lys Leu Gln Gln
1105                1110                1115                1120
Leu Met Glu Gln Ala Val Ala Pro Glu Ser Ser Glu Thr Leu Thr Asp
                1125                1130                1135
Ser Glu Thr Asn Gly Ser Thr Pro Leu Ala Asp Ser Asp Thr Ala Asp
            1140                1145                1150
Gly Thr Gln Gln Asp Glu Thr Ile Asp Ser Gln Asp Ser Lys Ala Thr
        1155                1160                1165
Ala Ala Val Arg Gln Ser Gln Val Thr Glu Glu Glu Ala Ala Thr Ala
    1170                1175                1180
Gln Lys Glu Glu Pro Ser Thr Leu Pro Asn Asn Val Pro Ala Gln Glu
1185                1190                1195                1200
Glu His Gly Glu Glu Pro Gly Arg Asp Val Leu Glu Pro Thr Gln Gln
            1205                1210                1215
Glu Leu Thr Ala Ala Ala Val Pro Val Leu Ala Lys Thr Glu Val Gly
        1220                1225                1230
Gln Glu Gly Glu Val Asp Trp Leu Asp Gly Glu Lys Val Lys Glu Glu
        1235                1240                1245
Gln Glu Val Phe Val His Ser Gly Pro Asn Ser Gln Lys Ala Ala Asp
    1250                1255                1260
Val Thr Tyr Asp Ser Glu Val Met Gly Val Ala Gly Cys Gln Glu Lys
1265                1270                1275                1280
Glu Ser Thr Glu Val Gln Ser Leu Ser Leu Glu Glu Gly Glu Met Glu
                1285                1290                1295
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Thr Asp Val Glu Lys Glu Lys Arg Glu Thr Lys Pro Glu Gln Val Ser
            1300                1305                1310
Glu Glu Gly Glu Gln Glu Thr Ala Ala Pro Glu His Glu Gly Thr Tyr
            1315                1320                1325
Gly Lys Pro Val Leu Thr Leu Asp Met Pro Ser Ser Glu Arg Gly Lys
        1330                1335                1340
Ala Leu Gly Ser Leu Gly Gly Ser Pro Ser Leu Pro Asp Gln Asp Lys
1345                1350                1355                1360
Ala Gly Cys Ile Glu Val Gln Val Gln Ser Leu Asp Thr Thr Val Thr
            1365                1370                1375
Gln Thr Ala Glu Ala Val Glu Lys Val Ile Glu Thr Val Val Ile Ser
            1380                1385                1390
Glu Thr Gly Glu Ser Pro Glu Cys Val Gly Ala His Leu Leu Pro Ala
            1395                1400                1405
Glu Lys Ser Ser Ala Thr Gly Gly His Trp Thr Leu Gln His Ala Glu
        1410                1415                1420
Asp Thr Val Pro Leu Gly Pro Glu Ser Gln Ala Glu Ser Ile Pro Ile
1425                1430                1435                1440
Ile Val Thr Pro Ala Pro Glu Ser Thr Leu His Pro Asp Leu Gln Gly
            1445                1450                1455
Glu Ile Ser Ala Ser Gln Arg Glu Arg Ser Glu Glu Glu Asp Lys Pro
        1460                1465                1470
Asp Ala Gly Pro Asp Ala Asp Gly Lys Glu Ser Thr Ala Ile Glu Lys
        1475                1480                1485
Val Leu Lys Ala Glu Pro Glu Ile Leu Glu Leu Glu Ser Lys Ser Asn
        1490                1495                1500
Lys Ile Val Leu Asn Val Ile Gln Thr Ala Val Asp Gln Phe Ala Arg
1505                1510                1515                1520
Thr Glu Thr Ala Pro Glu Thr His Ala Tyr Asp Ser Gln Thr Gln Val
            1525                1530                1535
Pro Ala Cys Arg Leu Asp Ser Arg Glu Pro Asn Arg Cys Trp Thr Lys
        1540                1545                1550
Met Lys Asp Ala Lys Met Lys His Pro Val Pro Gln Pro Arg Glu Asp
        1555                1560                1565
Leu Gln Val Leu Thr Val Leu Glu Ala Trp Ala Gln Pro Arg Lys Cys
    1570                1575                1580
Leu Pro Arg Leu Gln Leu Lys Ala Pro Val Ser Lys
1585                1590                1595
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<210> 5
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Myristylation site for src protein

<400> 5
Met Gly Ser Ser Lys Ser Lys Pro Lys Asp
 1   5   10

<210> 6
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Myristylation and palmitylation site for yes
     protein

<400> 6
Met Gly Cys Ile Lys Ser Lys Glu Asp Lys
 1   5   10

<210> 7
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Myristylation and/or palmitylation site for SSeCKS
     protein

<400> 7
Met Gly Ala Gly Ser Ser Thr Glu Gln Arg
 1   5   10

<210> 8
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Myristylation site for G protein

<400> 8
Met Gly Ala Gly Ala Ser Ala Glu Glu Lys
 1   5   10

<210> 9
<211> 10

<212> PRT
<213> Artificial Sequence

<220>
<223> Myristylation and palmitylation site for G protein

<400> 9
Met Gly Cys Thr Leu Ser Ala Glu Asp Lys
 1   5   10

<210> 10
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Palmitylation site for GAP-43 protein

<400> 10
Met Leu Cys Cys Met Arg Arg Thr Lys Gln
 1   5   10

<210> 11
<211> 23
<212> PRT
<213> Bos taurus

<400> 11
Lys Arg Phe Ser Ser Lys Lys Ser Phe Lys Leu Ser Gly Phe Ser Phe
 1   5   10   15
Lys Lys Asn Lys Lys Glu Ala
            20

<210> 12
<211> 23
<212> PRT
<213> Mus musculus

<400> 12
Lys Arg Phe Ser Ser Lys Lys Ser Phe Lys Leu Ser Gly Phe Ser Phe
 1   5   10   15
Lys Lys Ser Lys Lys Glu Ala
            20

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<210> 13
<211> 17
<212> PRT
<213> Unknown

<220>
<223> Phosphorylation consensus site for MacMARCKS/F52
      protein

<400> 13
Lys Lys Phe Ser Ser Lys Lys Pro Phe Lys Leu Ser Gly Phe Ser Phe
 1               5                  10                  15
Arg



<210> 14
<211> 20
<212> PRT
<213> Unknown

<220>
<223> Calmodulin binding domain for myosin light chain
      kinase protein

<400> 14
Lys Arg Arg Trp Lys Lys Ala Phe Ile Ala Val Ser Ala Ala Ala Arg
 1               5                  10                  15
Phe Lys Lys Cys
            20



<210> 15
<211> 23
<212> PRT
<213> Rattus norvegicus

<400> 15
Glu Thr Thr Ser Ser Phe Lys Lys Phe Phe Thr His Gly Thr Ser Phe
 1               5                  10                  15
Lys Lys Ser Lys Glu Asp Asp
            20



<210> 16
<211> 23
<212> PRT
<213> Rattus norvegicus
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<400> 16
Lys Leu Phe Ser Ser Ser Gly Leu Lys Lys Leu Ser Gly Lys Lys Gln
 1   5   10   15
Lys Gly Lys Arg Gly Gly Gly
            20

<210> 17
<211> 23
<212> PRT
<213> Rattus norvegicus

<400> 17
Glu Gly Ile Thr Pro Trp Ala Ser Phe Lys Lys Met Val Thr Pro Lys
 1   5   10   15
Lys Arg Val Arg Arg Pro Ser
            20

<210> 18
<211> 23
<212> PRT
<213> Rattus norvegicus

<400> 18
Glu Gly Val Ser Thr Trp Glu Ser Phe Lys Arg Leu Val Thr Pro Arg
 1   5   10   15
Lys Lys Ser Lys Ser Lys Leu
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<210> 19
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> SSeCKS phosphorylation consensus sequence

<221> VARIANT
<222> (3)...(3)
<223> valine or isoleucine

<221> VARIANT
<222> (4)...(5)
<223> any amino acid

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<221> VARIANT
<222> (7)...(7)
<223> any amino acid

<221> VARIANT
<222> (11)...(11)
<223> lysine or arginine

<221> VARIANT
<222> (12)...(12)
<223> any amino acid

<221> VARIANT
<222> (16)...(16)
<223> lysine or arginine

<221> VARIANT
<222> (18)...(18)
<223> lysine or arginine

<221> VARIANT
<222> (19)...(19)
<223> any amino acid

<221> VARIANT
<222> (20)...(20)
<223> lysine or arginine

<400> 19
Glu Gly Xaa Xaa Xaa Trp Xaa Ser Phe Lys Xaa Xaa Val Thr Pro Xaa
 1               5                   10                  15
Lys Xaa Xaa Xaa
          20


<210> 20
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> consensus sequence

<400> 20
Trp Ala Gly Trp Arg Lys Lys
 1                   5
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<210> 21
<211> 54
<212> PRT
<213> Notophthalmus viridescens

<220>
<221> VARIANT
<222> (20)...(46)
<223> any amino acid

<400> 21
Ser Pro Leu Lys Ser Pro Tyr Lys His Pro Glu Gly Leu Leu Ser Pro
 1   5   10   15
Thr Lys Met Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
            20   25   30
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Leu Ser
        35   40   45
Ser Ser Glu Arg Leu Arg
    50

<210> 22
<211> 14
<212> PRT
<213> Rattus norvegicus

<220>
<221> VARIANT
<222> (6)...(9)
<223> any amino acid

<400> 22
Lys Lys Leu Phe Ser Xaa Xaa Xaa Xaa Lys Lys Leu Ser Gly
 1   5   10

<210> 23
<211> 10
<212> PRT
<213> Rattus norvegicus

<400> 23
Met Gly Ala Gly Ser Ser Thr Glu Gln Arg
 1   5   10

<210> 24
<211> 26

<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

<400> 24
gtgactggtg aggcctcaac caagtc
26

<210> 25
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

<400> 25
gtgactggtg agtactcaac caagtc
26

<210> 26
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

<400> 26
ggaagtccct tgtcgagcct cttcagtagc
30

<210> 27
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

<400> 27
gctcaggctt aagctcgctg tctggg
26

<210> 28
<211> 24

<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

<400> 28
cccttgaaga aaagcttcag tagc
24

<210> 29
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

<400> 29
ggcttaaaga agtcgtctgg gaag
24

<210> 30
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

<400> 30
cccttgtcga gcagcttcag tagc
24

<210> 31
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

<400> 31
ggcttaagct cgtcgtctgg gaag
24

<210> 32
<211> 16

<212> PRT
<213> Artificial Sequence

<220>
<223> Penetratin peptide

<400> 32
Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys
 1   5   10   15

<210> 33
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> SSeCKS cyclin binding site

<400> 33
Leu Lys Lys Leu Phe Ser Ser Ser Gly Leu Lys Lys Leu Ser Gly Lys
 1   5   10   15

<210> 34
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> SSeCKS mutated cyclin binding site

<400> 34
Leu Ser Ser Ser Phe Ser Ser Ser Gly Leu Ser Ser Ser Ser Gly Lys
 1   5   10   15

<210> 35
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> SSeCKS mutated phosphoserine cyclin binding site

<221> VARIANT
<222> (6)...(6)
<223> phosphatidyl serine

<221> VARIANT
<222> (14)...(14)
<223> phosphatidyl serine

<400> 35
Leu Lys Lys Leu Phe Ser Ser Ser Gly Leu Lys Lys Leu Ser Gly Lys
 1   5   10   15